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C:Date: 15-Feb-1997 #sequence_revision 13-Mar-1997 #text_change C:Accession: S68691; S74313; S88683
R:Kjeldsen, L.; Cowland, J.B.; Johnsen, A.H.; Borregaard, N. FEBS Lett. 380, 246-250, 1996
A:Title: SGP28, a novel matrix glycoprotein in specific granules A:Reference number: S68691; MUID:96186934; PMID:8601434
A:Accession: S68691
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S68691
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A;Residues: 33-83;96-143;165-217;221-226 <KJL>
R;Kraetzschmar, J; Haendler, B; Eberspaecher,
Eur J. Blochem. 236, 827-836, 1996
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A; Residues: 1-245 < KJE>
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210	211.5	214	217	218	220.5	222	223.5	224	224.5	226.5	227.5	227.5	228.5	234	234.5	
14.6	14.7	14.9	15.1	15.2	15.4	15.5	15.6	15.6	15.6	15.8	15.8	15.8	15.9	16.3	16.3	
175	205	177	212	205	164	205	164	207	164	211	207	164	164	212	209	
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S43894	T48294	T08126	T22437	F44583	S52627	A37329	S37166	T31959	T04299	T19859	T22436	S39474	S52626	В37330	T19848	
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ALIGNMENTS

ovel matrix glycoprotein in specific granules S68691; MUID:96186934; PMID:8601434

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human neutrophil

17-Nov-2000

EMBL: X94323; NID: g1213612; PIDN: CAA63984.1;

PID:g1213613

B.; Eberspaecher, U.; Roosterman, 1996

D.; Donner, P.; Schl

A:Molecule type: mRNA A:Residues: 1-105, 'S',107-245 <KRA> A:Residues: 1-105, 'S',107-240; NID:gl262818; PIDN:CAA64527.1; PID:gl262819 A:Cross references: EMBL:X95240; NID:gl262818; PIDN:CAA64527.1; PID:gl262819 C:Genetics: A:Gene: SGP28 C:Superfamily: cysteine-rich secretory protein 1 C:Superfamily: cysteine-rich secretory protein 1 F:1-19/Domain: signal sequence #status predicted <SIG> F:20-245/Product: neutrophil granules matrix glycoprotein SGP28 #status predicted A;Title: The human cysteine-rich secretory protein (CRISP) family. Primary structure A;Reference number: S68681; MUID:96270732; PMID:8665901 A;Accession: S68683 A;Status: preliminary 밁 В Ş В Š Best Loc Matches Query Match 194 181 121 134 61 74 14 Local LYVPYEQGAPCASCPDNCDDGLCTNGCKYEDLYSNCKSLKLTLTCKHQLVRDSCKASCNC LKMEWNKEAAANAQKWANQCNYRHSNPKDRMTSLKCGENLYMSSAPSSWSQAIQSWFDEY LKMEWNKEAAANAQKWANQCNYRHSNPKDRMTSLKCGENLYMSSAPSSWSQAIQSWFDEY Similarity Conservative 95.5%; Score 1371; DB 2; 100.0%; Pred. No. 1.7e-102; Live 0; Mismatches 0; Length 245; Indels 0; Gaps 193 120 60 253 180 0 ₹

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testicular protein Tpx-1 - rat
C;Species: Rattus norvegicus (Norway rat)
C;Date: 21-Aug-1998 #sequence_revision 21
C;Accession: JE0204
R; Maeda, T.; Sakashita, M.; Ohba, Y.; Nakanishi, Y. Biochem. Biophys. Res. Commun. 248, 140-146, 1998 A;Ttle: Molecular cloning of the rat tpx-1 responsible A;Reference number: JE0204; MUID:98340864; PMID:9675100
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A;Title: Cloning and mapping of a testis-specific A;Reference number: A33329; MUID:90129048; PMID:26 A;Accession: B33329
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cysteine-rich secretory protein 2 type I precursor - human N;Alternate names: testis-specific protein C;Species: Homo saplens (man) C;Date: 09-Mar:1990 #sequence_revision 09-Mar-1990 #text_change
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A; Residues: 1-243 < KRA>
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A; Residues: 1-243 <KAS>
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Pred. No. 7.6e-72
5; Mismatches 4:
                                                                       21-Aug-1998 #text_change
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C;Comment:
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C;Species: Mus musculus (house mouse)
C;Date: 09-Mar-1990 #sequence_revision
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C; Superfamily:
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A; Residues: 1-243 <MAE>
A; Cross-references: DDB
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59.6%;
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Pred No. 3.3e-59;
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Pred. No. 5.8e-61;
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as a cell adhesion protein for 1
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RESULT 5
A49202
A49202
Cysteine-rich secretory protein-1 - mouse
N;Alternate names: CRISP-1
C;Species: Mus musculus (house mouse)
C;Date: 19-Dec-1993 #sequence_revision 18-Nov-1994 #text_change 29-Sep-1999
C;Accession: A49202
C;Accession: A49202
R;Haendler, B.; Kratzschmar, J.; Theuring, F.; Schleuning, W.D.
Endocrinology 133, 192-198, 1993
A;Title: Transcripts for cysteine-rich secretory protein-1 (CRISP-1; DE/AEG) and A;Reference number: A49202; MUID:93307144; PMID:8319566
A;Accession: A49202
A;Status: preliminary
A;Molecule type: nucleic acid
A;Residues: 1-244 <HAED>
A;Cross-references: GB:L05559; NID:9309190; PIDN:AAA37460.1; PID:9309191
A;Experimental source: NMRI, epididymis, salivary gland
A;Note: sequence extracted from NCBI backbone (NCBIN:134675, NCBIP:134676)
C;Superfamily: cysteine-rich secretory protein 1
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                                                                                                     A; Molecule type: mRNA
A; Residues: 1-246 <CHA>
A; Residues: 1-246 <CHA>
A; Cross references: GB.M31173; NID:g202772; PIDN:AAB59716.1; PID:g202773
A; Brooks, D.E.; Means, A.R.; Wright, E.J.; Singh, S.P.; Tiver, K.K.
Bur. J. Biochem. 161, 13-18, 1986
Bur. J. Biochem. 161, 13-18, 1986
A; Title: Molecular cloning of the cDNA for androgen-dependent sperm-coat:
A; Reference number: A24609; MUID:87053995; PMID:3780731
                                                                                                                                                                                                                                                                         R;Charest, N.J.; Joseph, D.R.; Wilson, E.M.; French, F.S. Mol. Endocrinol. 2, 999-1004, 1988
A;Title: Molecular cloning of complementary deoxyribonucleic A;Reference number: A40918; MUID:89039913; PMID:2460753
A;Accession: A40918
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A24609
A244609
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                                           A; Residues: 1-246 <BRO>
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Cross-references: GB:X04643; NID:g56112; PIDN:CAA28304.1; Superfamily: cysteine-rich secretory protein 1
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Pred. No. 4e-54; 
9; Mismatches 7
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121 KGLIFGVGPKQNVSVVGHHTQVVWKSNLQVACGVAECP-ENPLRYFYVCRYCPVLNYSGH

NDEDEGVGPKTPNAVVGHYTQVVWYSSYLVGCGNAYCPNQKVLKYYYVCQYCP----AGN

120 189

LIMEWNYDAQVIAQQRADKCTFSHSPIELRTTNLKCGENLFMSSYLVPWSSVIQGWYNES

WANRLYVPYEQGAPCASCPDNCDDGLCTNGCKYEDLYSNCKSLKLTLTCKHQLVRDSCKA

YPSRPYLAYTARAPCASCPDRCEDGLCTKSCQYKDMSFWCK--RLEYVCKHPGLKKRCLA

179 249 237 8

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SCNC

250 238

TCQC

253

80 8

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190 180

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cysteine-rich secretory protein-3 - mouse
N;Alternate names: CRISP-3
C;Species: Mus musculus (house mouse)
C;Date: 19-Dec-1993 #sequence_revision 18-Nov-1994 #text_change
C;Accession: B49202
C;Accession: B4
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F;1-19/Domain: signal sequence #status predicted <SIG>
F;20-246/Product: acidic epididymal glycoprotein #stat
                                                                                                                                                                                                                                                                                                                                                    A;Cross-references: GB:L05560: NID:g309192; PIDN:AAA37461.1; PID:g309193 A;Experimental source: NMRI, epididymis, salivary gland A;Note: sequence extracted from NCBI backbone (NCBIN:134677, NCBIP:134670 C;Superfamily: cysteine-rich secretory protein 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Title: Transcripts for cysteine-rich secretory protein-1 (CRISP-1; A;Reference number: A49202; MUID:93307144; PMID:8319566 A;Accession: B49202
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Endocrinology 133, 192-198, 1993
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A; Residues: 1-241 <HAE>
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                                                                                            MALMLVLFFLAAVLPPSLLQDNSQENSLEKLSTSKKSVQEEIVSKHNQLRRKVSPSGSDL
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LKMEWNKEAAANAQKWANQCNYRHSNPKDRWTSLKCGENLYMSSAPSSWSQAIQSWFDEY
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118; Conservat
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                                                                                                                                                                                                                        Score 617.5; DB 2;
Pred. No. 3.3e-42;
10; Mismatches 79;
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Cysteine-rich secretory protein 1 precursor - human
C; Species: Homo sapiens (man)
C; Date: 15-Feb-1997 *sequence_revision 13-Mar-1997 *text_change 20-:
C; Date: 15-Feb-1997 *sequence_revision 13-Mar-1997 *text_change 20-:
C; Accession: S66884
R; Kraetzschmar, J.; Haendler, B.; Eberspaecher, U.; Roosterman, D.;
Eur. J. Blochem. 236, 827-836, 1996
A; Title: The human cysteine-rich secretory protein (CRISP) family. I
A; Reference number: S68681; MUID:96270732; PMID:8665901
A; Reference number: S68681; MUID:96270732; PMID:8665901
A; Rocession: S68684
A; Status: prellminary
A; Molecule type: mRNA
A; Residues: 1-249 <KRA>
A; Residues: 1-249 <KRA>
A; Cross-references: EMEL:X95237; NID:g1262814; PIDN:CAA64524.1; PID
C; Superfamily: cysteine-rich secretory protein 1
F;1-21/Domain: signal sequence *status predicted <SIG>
F;22-249/Product: cysteine-rich secretory protein 1 *status predicted
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                                                                                                                                                     C;Superfamily: cysteine-rich secretory protein 1
F;1-21/Domain: signal sequence #status predicted <SIG>
F;22-178/Product: cysteine-rich secretory protein 1-delta #status experimental <MAT>
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                                                                                                                                                                                                                                                                                                                                             A; Reference number: A; Accession: S68681
                                                                                                                                                                                                                                                                                                                                                                A:Title: The human cysteine-rich secretory A:Reference number: S68681; MUID:96270732;
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C;Date: 15-Feb-1997 #sequence_revision 13-Mar-1997 #
C;Accession: S68681; S74302
R;Kraetzschmar, J.; Haendler, B.; Eberspaecher, U.;
Eur. J. Blochem. 236, 827-836, 1996
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A; Residues: 1-178 <KRA>
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Matches 111
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Best Local :
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                       LLELVAG--LLP--SFPANEDKDPAFTALLTTQTQVQREIVNKHNELRRAVSPPARNMLK 75
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LYVPYEQGAPCASCPDNCDDGLCTNGCKYEDLYSNCKSLKLTLTCKHQLVRDSCKASCNC
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  LLFLVAAACLLPMLSMKKKSARD-QFNKLVTDLPNVQEEIVNIHNALRRRVVPPASNMLK
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45.5%; Pro
ative 26;
                                                                                                   26.0%;
                                                                                                                                                                                                                                                                                    NID: g1262812;
                                                                                                     Score 373.5; DB 2; Pred. No. 8.1e-23;
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Pred. No. 3.8e-35;
6; Mismatches 100
                                                                                                                                                                                                                                                                                                                                                                protein (CRISP)
PMID:8665901
                                                                                                                                                                                                                                                                                    PIDN: CAA64525.1;
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                                                                                                               R:Murphy, E.V.; Zhang, Y.; Zhu, W.; Biggs, J. Gene 159, 131-135, 1939
A;Title: The human glioma pathogenesis-related protein A;Reference number: JC4131; MUID:95331646; PMID:7607567
A;Accession: JC4131
                                                                                                                                                                                                                                      glioma pathogenesis-related protein - }
C;Species: Homo sapiens (man)
C;Date: 02-Aug-1995 #sequence_revision
                                      A; Experimental source:
                                                    A;Residues: 1-219 <MUR>
A;Cross references: GB:U16307; NID:gl100927; PIDN:AAA82731.1;
                                                                              A; Molecule type: mRNA
A; Residues: 1-219 < MU
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C; Genetics: A; Gene: GDE
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A, Gene: rtvp-1
A, Gene: rtvp-1
C, Superfamily: yellowjacket venom allergen antigen
C, Superfamily: yellowjacket venom allergen antigen
F, 1-21/Domain: signal sequence #status predicted
F, 22-266/Product: testis-specific, vespid, and pail
F, 22
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F;233-255/Domain: transmembrane #status
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A; Molecule type: protein
A; Residues: 97-100;114-120;134-144 <RIC2>
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A; Residues: 1-266 <RIC1>
A; Cross-references: EMBL:X91911; NID:g1030052;
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R; Rich, T.; Chen, P.; Furman,
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      SGFDALSNGAHFICNYGPGGNYPT --
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Pred. No. 5.3e-19;
      -WPYKRGATCSACPNNDKCLDNLCVN
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predicted <TMM>
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venom allergen antigen Vesp c 5.01 - European hornet
C;Species: Vespa crabro (European hornet)
C;Date: 27-Jun-1994 #sequence_revision 27-Jun-1994 #text_change 11-Jan-2000
C;Accession: G44583; G44522
R;Hoffman, D.R.
J. Allergy Clin. Immunol. 92, 707-716, 1993
A;Hitle: Allergens in hymenoptera venom XXV: the amino acid sequences of ant A;Reference number: A44583; MUID:94044316; PMID:8227862
A;Accession: G44583
A;Status: preliminary
A;Residues: 1-202 <HOFP
C;Superfamily: yellowjacket venom allergen antigen 5
                                          venom allergen antigen Vesp c 5.02 - European hornet C; Species: Vespa crabro (European hornet) C; Cpate: 27-Jun-1994 *sequence_revision 27-Jun-1994 *text_cha C; Accession: H44583; H44522 R; Hoffman, D.R. J. Allergy Clin. Immunol. 92, 707-716, 1993 A; Title: Allergens in hymenoptera venom XXV: the amino acid A; Reference number: A44583; MUID:94044316; PMID:8227862 A; Accession: H44583 A; Status: preliminary
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C:Superfamily: yellowjacket venom
C;Keywords: brain
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                 Molecule type: protein Residues: 1-202 <HOF>
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                                                                                                                                                                                                                                                                                                              149 VGHYTQVVWYSSYLVGCGNAYCPNQKVLKYYYVCQYCPAGNWAN 192
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                                                                                                                                                                                                                                                                                                                                                                 CNYRHSNPKDRMTSLKCGENLYMSSAPS----SWSQAIQSWFDEYNDFDFGVGPKTPNAV 148
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AGLLPSFPANEDKDPAFTALLTTQTQVQREIVNKHNELRRAVSPPARNMLKMEWNKEAAA 84
                                                                                                                                                                                                                                                                                                                                                                                                              LTKQENL--EILKQHNEFRQKVARGLETRGNPGPQPPAKSMNTLVWNDELAQIAQVWANQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IAKAWASNCQFSHNTRLKPPHKLHPNFTSL--GENIWTGSVPIFSVSSAITNWYDEIQDY
                                                                                                                                                                                                                                                                              VGHYTQMVWAKTKEIGCGSIKYIENGWHRHYLVCNYGPAGNVGN
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yellowjacket venom
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Pred. No. 2.1e-14;
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R;Hoffman, D.R.

J. Allergy Clin. Immunol. 10

A;Title: Reactions to less c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Cross-references: GB:J03601; NID:g156714; PID:g156715 C;Superfamily: yellowjacket venom allergen antigen 5 F;1-23/Domain: signal sequence #status predicted <MAT> F;24-227/Product: antigen 5-2 #status predicted <MAT>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C;Accession: A31085
R;Fang, K.S.Y.; Vitale, M.; Fehlner, P.; King, T.P.
Proc. Natl. Acad. Sci. U.S.A. 85, 895-899, 1988
A;Title: cDNA cloning and primary structure of a white-face
A;Reference number: A94213; MUID:88124947; PMID:3422469
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  antigen 5-2 precursor - bald-faced hornet
C;Species: Vespula maculata (bald_faced hornet)
C;Date: 31-Mar-1990 *sequence_revision 31-Mar-1990 *text_change 11-Jan-2000
                                                                                   A;Molecule type: protein
A;Residues: 1-211 <HOF>
C;Superfamily: yellowjacket venom
C;Keywords: venom
                                                                                                                                                                                                                                                                                                 venom allergen Sol r 3 - black imported fire ant
C;Species: Solenopsis richteri (black imported fire ant)
C;Date: 06-Nov-1998 #sequence_revision 06-Nov-1998 #text,
C;Accession: B58853
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A; Residues: 1-227 <FAN>
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                                                                                                                                                                     A; Status: preliminary
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Matches 60
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100 PKDRWTSLKCGENLYMSSAPSSW----SQAIQSWFDEYNDFDFGVGPKTPN-AVVGHYTQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            64 KNEILKRHNDFRONVAKGLETRGKPGPOPPAKNMNVLVWNDELAKIAOTWANOCDFNHDD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      52 QREIVNKHNELRRAVS-------PPARNMLKMEWNKEAAANAQKWANQCNYRHSN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      VGHYTQMVWAKTKEIGCGSEDYIEDGWHRHYLVCNYGPAGNVGN: 195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                VGHYTQVVWYSSYLVGCGNAYCPNQKVLKYYYVCQYCPAGNWAN 192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CNYGHDNCRN-SAKYSVGQNIAEGSTSADNFVNVSNMVKMWEDEVKDYQYG-SPKNKLNK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LTKQENL -- EILKQHNEFRQKVARGLETRGNPGPQPPAKSMNTLVWNDELAQIAQVWANQ
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60; Conserv
ch 17.9%;
l Similarity 36.2%;
59; Conservative
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                                                                                                                                                                                                                                                           100, 679-683, 1997
                                                                                                                                                                                                                 common species of fire ants. MUID:98049167: PMID:9389299
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 265; DB 2;
Pred. No. 4.4e-14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 259; DB 2;
Pred. No. 1.5e-13;
1; Mismatches 55
    Score 257.5; DB 2;
Pred. No. 1.8e-13;
4; Mismatches 57;
                                                                                                            allergen
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Local

24;

Indels

23;

Gaps

6,

Sear	Db	Qy	Db	Qy	р	Qy
Search completed: March 14, 2003, 05:41:32	158 MHVGHYTQIVWAKTKKIGCGRIMFKEDNWNKHYLVCNYGPAGN 200	149 VGHYTQVVWYSSYLVGCGNAYCPNQKVLKYYYYCQYCPAGN 189	99 EHDACRN-VERFAVGONIAATSSSGKNKSTLSDMILLWYNEVKDFDNRWISSFPSDGNIL 157	96 RHSNPKDRMTSLKCGENLYMSSAPSSWSQAIQSWFDEYNDPDFGVGPKTPNAV 148	39 TDAEKDAIVNKHNELRORVASGKEMRGTNGPOPPAVKMPNLTWDPELATIAORWANOCTF 98	